

CM-1 V_H:

```

agg tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc agt agc tat ggc atg cac      60
Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr Gly Met His      20
1      5      10      15      20

tgg gtc cgc cag gct cca ggc aag ggg ctg gag tgg gtg gca gtt ata tca tat gat gga      120
Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ala Val Ile Ser Tyr Asp Gly      40
25      30      35      40

agt aat aaa tac tat gca gac tcc gtg aag ggc cga ttc acc atc tcc aga gac aat tcc      180
Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser      60
45      50      55      60

aag aac acg ctg tat ctg caa atg aac agc ctg aga gct gag gac acg gct gtg tat tac      240
Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr      80
65      70      75      80

tgt gcg aaa gac cgg tct tcg ggc tac tac ggt atg gac gtc tgg ggc caa ggc acc ctg      300
Cys Ala Lys Asp Arg Ser Ser Gly Tyr Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Leu      100
85      90      95      100

gtc acc
Val Thr      306

```

FIG. 1

CM-1 VL:

60	tcc tat gtg ctg act cag cca ccc tcg gtg tca gtg tcc cca gga cag acg gcc agg atc	15	gga cag acg gcc agg atc
	Ser Tyr Val Leu Thr Gln Pro Pro Ser Val Ser Val Ser Val Ser Val Ser Val Ser Val	20	Gly Gln Thr Ala Arg Ile
120	acc tgc tct gga gat gca ttg cca aag caa tat gct tat tgg tac cag cag aag cca ggc	35	gga cag acg gcc agg atc
	Thr Cys Ser Ser Gly Asp Ala Leu Pro Lys Gln Tyr Ala Tyr Trp Tyr Gln Gln Lys Pro Gly	40	Gly Gln Thr Ala Arg Ile
180	cag gcc cct gtg ctg gtg ata tat aaa gac agt gag agg ccc tca ggg atc cct gag cga	55	gga cag acg gcc agg atc
	Gln Ala Pro Val Leu Val Ile Tyr Lys Asp Ser Glu Arg Pro Ser Ser Ser Ser Ser Ser	60	Gly Gln Thr Ala Arg Ile
240	ttc tct ggc tcc agc tca ggg aca aca gtc acg ttg acc atc agt gga gtc cag gca gaa	75	gga cag acg gcc agg atc
	Phe Ser Gly Ser Ser Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Ser Ser Ser Ser Ser	80	Gly Gln Thr Ala Arg Ile
300	gac gag gct gac tat tac tgt caa tca gca gac agc agt ggt act tat gtg gta ttc ggc	95	gga cag acg gcc agg atc
	Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Ser Ser Ser Ser Ser Ser	100	Gly Gln Thr Ala Arg Ile
327	gga ggg acc aag ctg acc gtc cta ggt		
	Gly Gly Thr Lys Leu Thr Val Leu Gly		

FIG. 2



FIG. 3C



FIG. 3B



FIG. 3A



FIG. 4C

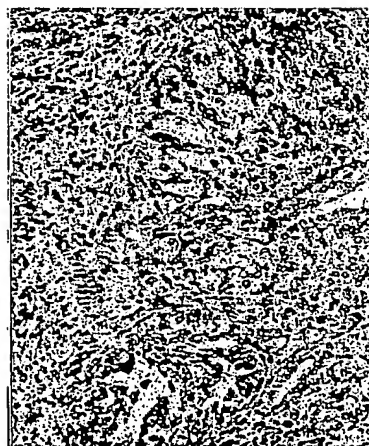


FIG. 4B



FIG. 4A



FIG. 5C



FIG. 5B



FIG. 5A

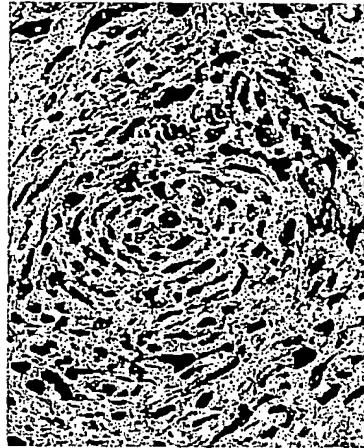


FIG. 6C

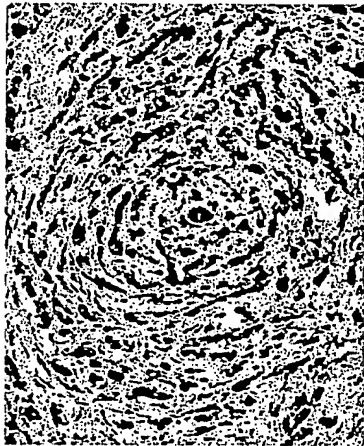


FIG. 6B

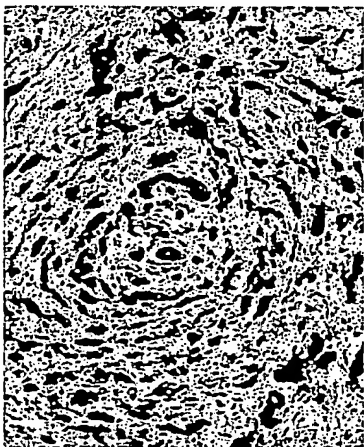


FIG. 6A



FIG. 7C

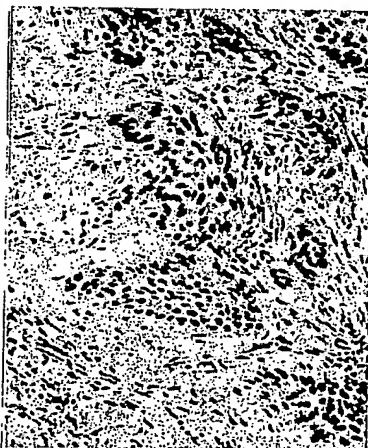


FIG. 7B



FIG. 7A

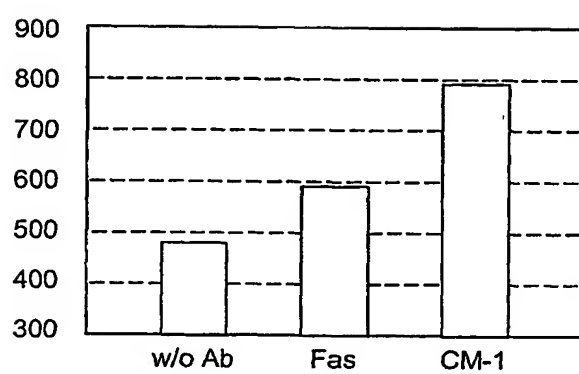


FIG. 8

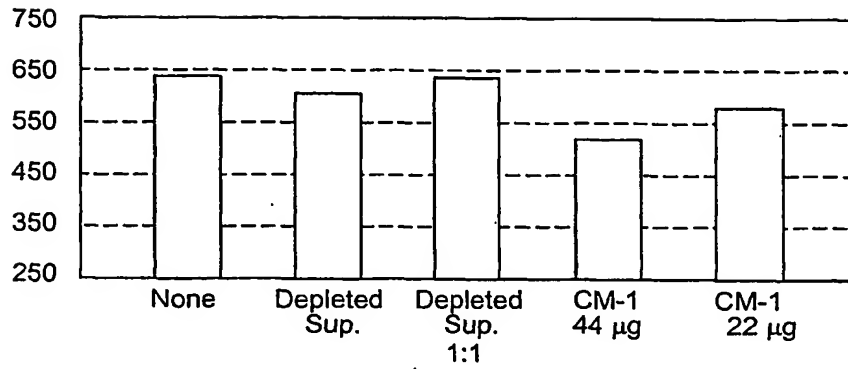


FIG. 9A

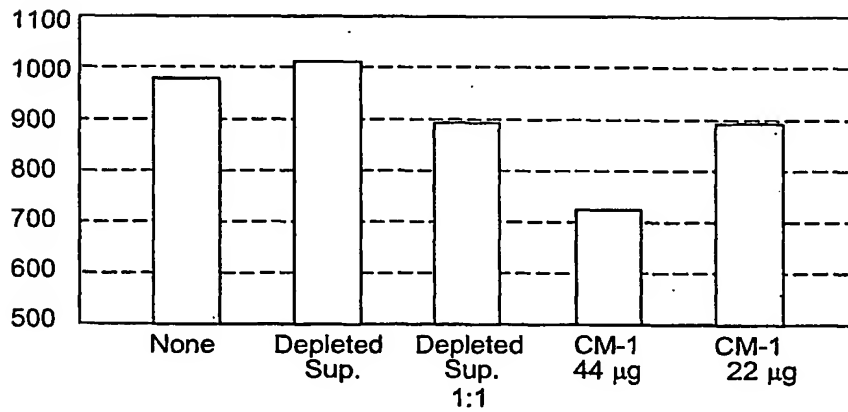


FIG. 9B